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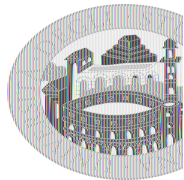
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# Investigating Diffusion-MRI based neurite density estimation model dependency: an in-vivo study on the HCP dataset



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**ISMRM 2018 Paris**  
**Time: 09:15, 18<sup>th</sup> June**  
**Number: 3235**  
**Computer N°: 41**

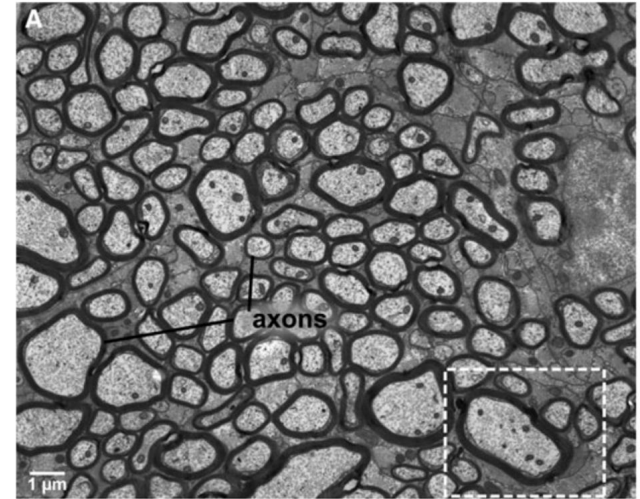


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Sherbrooke Connectivity Imaging Lab

# Neurite Density and Diffusion MRI

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- **Neurite density**<sup>1,2,3,4,5</sup> is one of the most promising microstructural features that can be estimated from **Diffusion-MRI** multi-shell data
- Recent years have seen a proliferation of **Multi-Compartment models** developed to estimate the neurite density



Nilsson et. al. "The role of tissue microstructure and water exchange in biophysical modelling of diffusion in white matter"  
*Magn Reson Mater Phy* (2013) 26:345370

# Multi-Compartment models

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- Multi-Compartment models represents the diffusion signal as a weighted sum of *compartments*

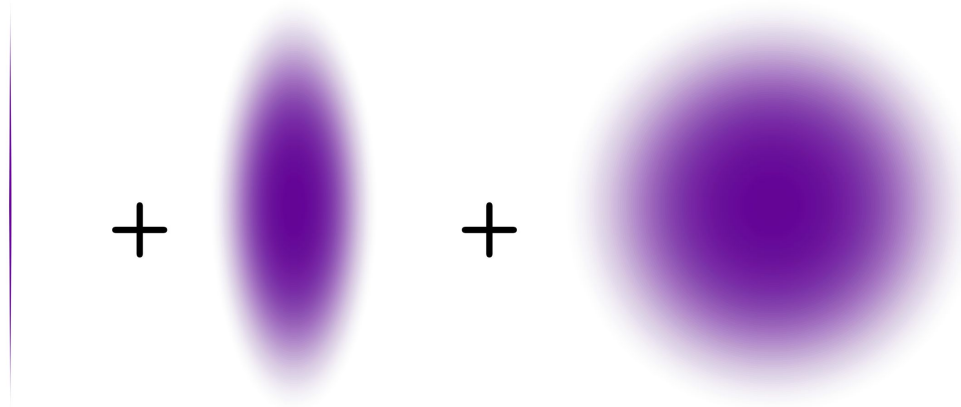
INTRA

EXTRA

CSF

+

+



# Neurite Density and Diffusion MRI

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- Neurite density can be calculated from Multi-Compartment models as the **intra-axonal volume fraction** ( $\nu_{ia}$ )

$$F(b, \vec{\mathbf{u}}, \vec{\mathbf{v}}) = \nu_{ia} F_{ia}(b, \vec{\mathbf{u}}, \vec{\mathbf{v}}) + \nu_{ea} F_{ea}(b, \vec{\mathbf{u}}, \vec{\mathbf{v}}) + \nu_{csf} F_{csf}(b)$$

- Recent years have seen a **proliferation**<sup>1,2,3,4,5,6,7,9,10</sup> of Multi-Compartment models
  - Each of these models makes different **assumptions** about the values of the diffusivity coefficient and the number of compartments
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# Spherical Mean Technique

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- This MC representation of the diffusion signal is valid only for fibers **aligned** in a single direction
- We can **convolve** the single fiber signal to the *fiber Orientation Distribution Function (fODF)*

$$E(b, \vec{u}) = \int_{\vec{v} \in S^2} \rho(\vec{v}) F(b, \vec{u}, \vec{v}) d\vec{v}$$

$$\rho(\vec{v}) = \sum_{l=0, \text{even}}^{\infty} \sum_{m=-l}^l c_{lm} Y_l^m(\vec{v})$$



$$K_l(b) Y_l^m(\vec{u}) = \int_{\vec{v} \in S^2} F(b, \vec{u}, \vec{v}) Y_l^m(\vec{v}) d\vec{v}$$


$$E(b, \vec{u}) = \sum_{l=0, \text{even}}^{\infty} \sum_{m=-l}^l c_{lm} K_l(b) Y_l^m(\vec{u})$$

- With  $Y_l^m(\vec{v})$  are the real **Spherical Harmonics** (SH) functions

# Spherical Mean Technique

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$$E(b, \vec{\mathbf{u}}) = \sum_{l=0, \text{even}}^{\infty} \sum_{m=-l}^l c_{lm} K_l(b) Y_l^m(\vec{\mathbf{u}})$$


$$\begin{aligned} \bar{E}(b) &= \frac{1}{4\pi} \int_{\vec{\mathbf{u}} \in \mathcal{S}^2} E(b, \vec{\mathbf{u}}) d\vec{\mathbf{u}} \\ &= \frac{1}{4\pi} K_0(b) \end{aligned}$$

- The **mean** of the signal depends **only** on the **microstructural kernel** and not on the **fiber orientation**<sup>7,9,10</sup>
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# Aims

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In this work:

- We will compare the neurite density estimated using **three** different Multi-Compartment models
  - We evaluate its **inter-subject reproducibility**
  - We evaluate the effect of the other model **parameters** on its estimation in-vivo
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# Multi-Compartment models

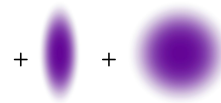
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## NODDI-SH

$$E(\mathbf{b}) = (1 - \nu_{csf})(E_{ia}(\mathbf{b}, \lambda_{\parallel}^{ia}) + \nu_{ea}E_{ea}(\mathbf{b}, \lambda_{\parallel}^{ea}, \lambda_{\perp}^{ea})) + \nu_{csf}E_{csf}(b)$$

$$\lambda_{\parallel}^{ia} = \lambda_{\parallel}^{ea} = 1.7 \cdot 10^{-3} \text{ mm}^2/\text{s}$$

$$\lambda_{\perp}^{ea} = \lambda_{\parallel}^{ea} \frac{\nu_{ea}}{\nu_{ia} + \nu_{ea}}$$



## BS-SH

$$E(\mathbf{b}) = \nu_{ia}E_{ia}(\mathbf{b}, \lambda_{\parallel}^{ia}) + \nu_{ea}E_{ea}(\mathbf{b}, \lambda_{\parallel}^{ea}, \lambda_{\perp}^{ea})$$

$$\lambda_{\parallel}^{ia} = 1.7 \cdot 10^{-3} \text{ mm}^2/\text{s}$$

$$\lambda_{\parallel}^{ea} = \lambda_{\perp}^{ea}$$



## MC-MDI

$$E(\mathbf{b}) = \nu_{ia}E_{ia}(\mathbf{b}, \lambda_{\parallel}^{ia}) + \nu_{ea}E_{ea}(\mathbf{b}, \lambda_{\parallel}^{ea}, \lambda_{\perp}^{ea})$$

$$\lambda_{\parallel}^{ia} = \lambda_{\parallel}^{ea}$$

$$\lambda_{\perp}^{ea} = \lambda_{\parallel}^{ea}(1 - \nu_{ia})$$



# Human Connectome Project (HCP)

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- We considered 10 subjects of the **Human Connectome Project**<sup>8</sup>
- b-values = [1000, 2000, 3000] s/mm<sup>2</sup>
- 90 gradients per shell plus 18 b 0
- $\Delta = 43.1\text{ms}$  and  $\delta = 10.6\text{ms}$

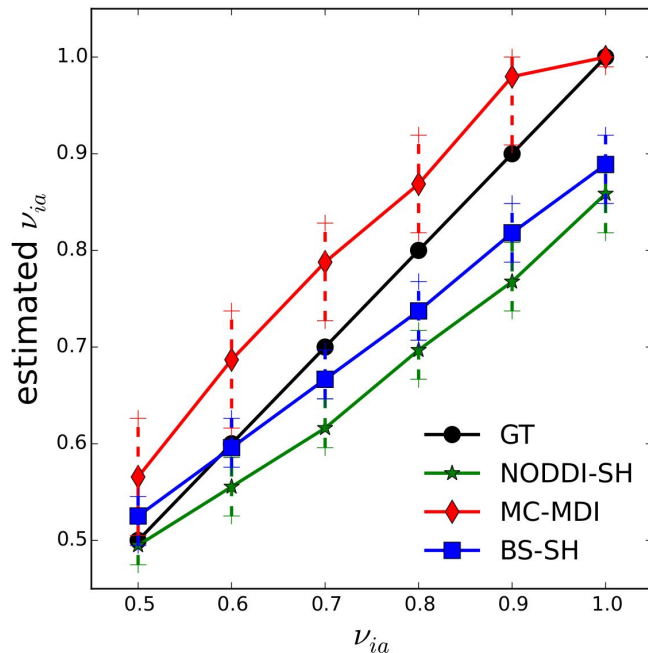


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# Synthetic data Results (from MICCAI 2017)

Our previous results<sup>7</sup> on **synthetic data** shown that:

- NODDI-SH and BS-SH tend to **underestimate** the neurite density
- MC-MDI tends to **overestimate** it

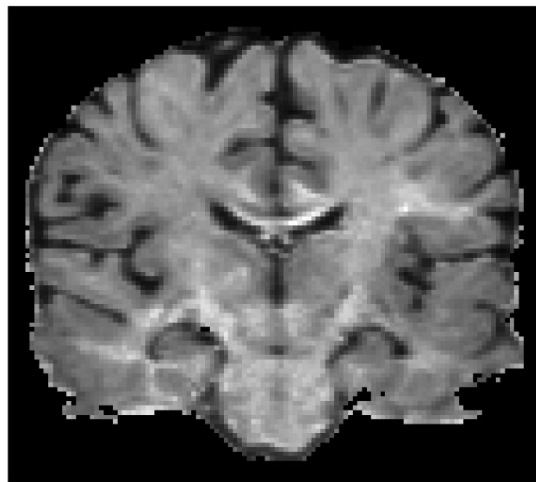


Zucchelli, M., Descoteaux, M., Menegaz, G. (2017). Proceedings of MICCAI, Workshop on Computational Diffusion MRI (CDMRI)", Canada.

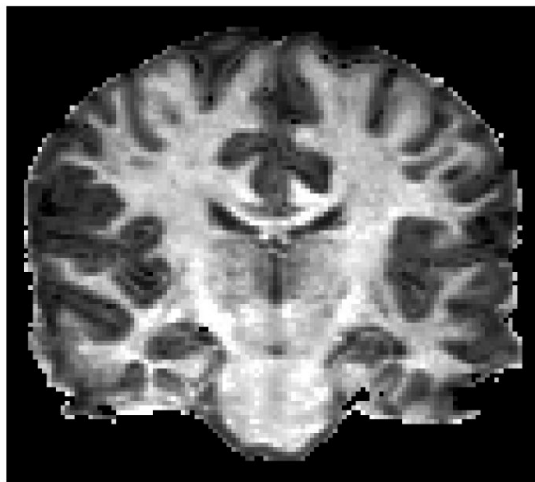
# HCP Results

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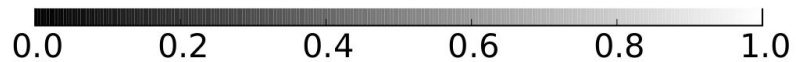
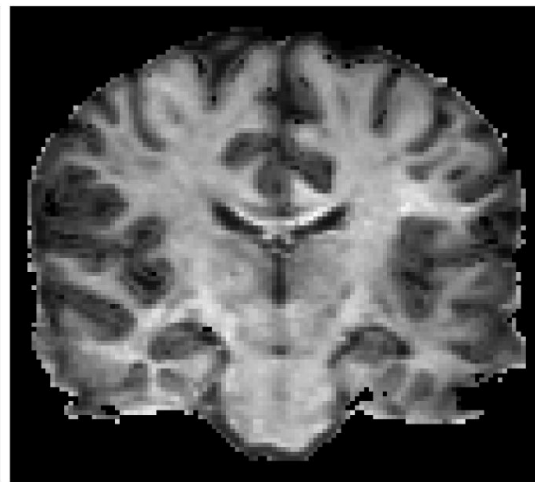
NODDI-SH



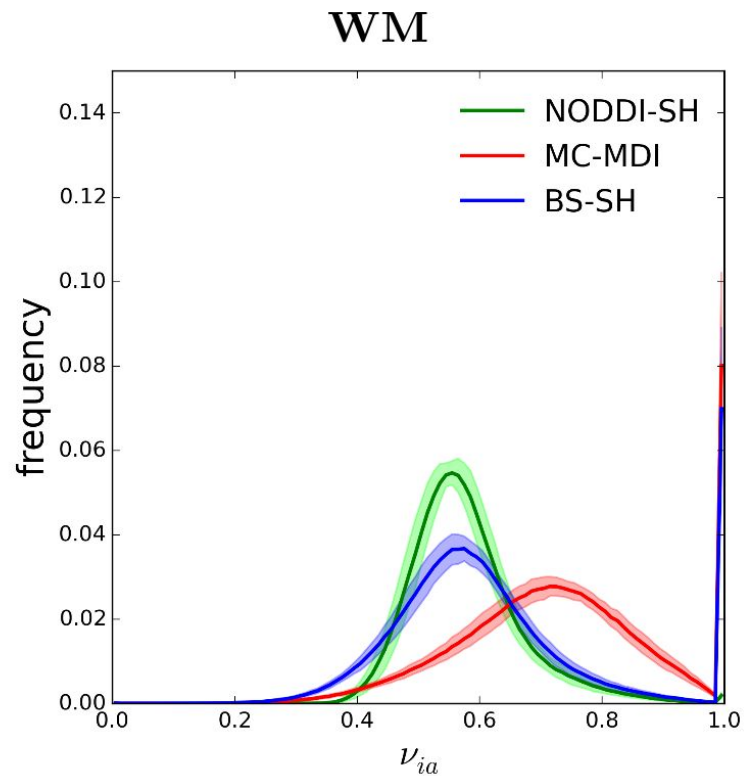
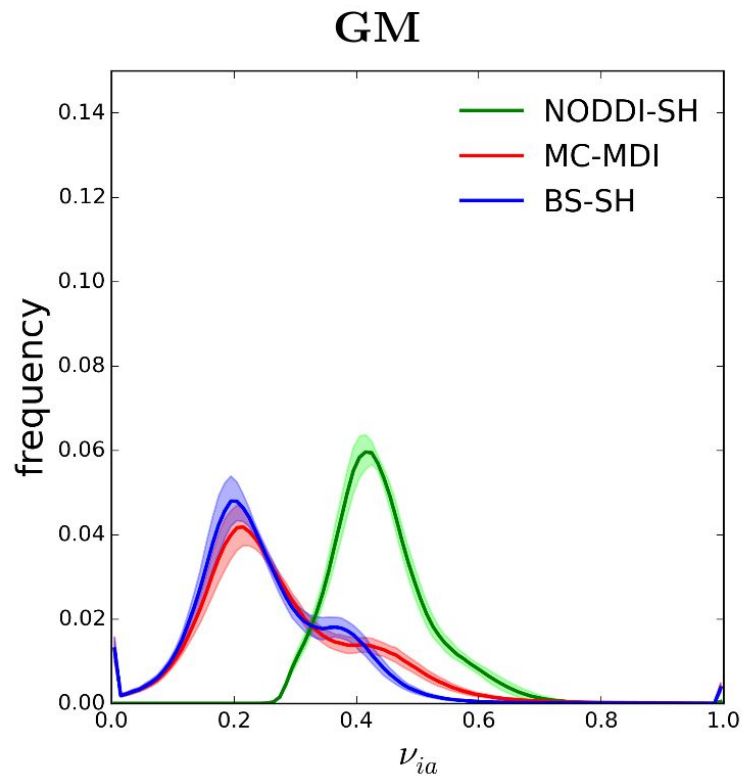
MC-MDI



BS-SH



# HCP Results



# Conclusions

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- **Neurite density** has a well defined numerical range and it is **stable** across healthy subjects
  - However, its values strongly depend on the **choice of the model** used to calculate it
  - Our results suggest that it could potentially be used as a **feature** to discriminate between healthy brains and pathological conditions
  - However, it is extremely important to keep in mind that its values are only proportional to the **real underlying neural density** and to compare it only with studies that use exactly the same model for its estimation
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